

SEQUENCE LISTING

<110> TORDO, NOEL
PERRIN, PIERRE
JACOB, YVES
BAHLOUL, CHOKRI
<120> CHIMERIC LYSSAV

8AHLOUL, CHORRI
<120> CHIMERIC LYSSAVIRUS NUCLEIC ACIDS AND POLYPEPTIDES
<130> 03495-0188-00000
<140> 09/549,519
<141> 2000-04-14
<150> 60/129,501
<151> 1999-04-15
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<170> PatentIn Ver. 2.1

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<213> Artificial Sequence

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<223> Description of Artificial Sequence: Primer

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ctagagee accatggite eteaggetet eetg

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<212> DNA

<213> Artificial Sequence

<220×

<223> Description of Artificial Sequence: Primer

<400> 2 attgatcaac tgaccgggag ggc

23

34

<210> 3

<211> 98

<212> DNA

<213> Artificial Sequence

<220>

<400> 3
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ccattgtgtt ttgggaagaa tteeceeec ggteagtt 98

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<210> 4
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aaacaggaga gcctgaggaa ccatggtggc ggctctag
<210> 5
<211> 54
<212> DNA
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<223> Description of Artificial Sequence: Synthetic
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aatttcccaa tctacaccat cccggataaa atcggaccgt ggtcacctat tccg
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<212> DNA
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aattoggaat aggtgaccac ggtccgattt tatccgggat ggtgtagatt ggga
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<223> Description of Artificial Sequence: Primer
ccgtggtcac ctattgatat aaaccatctc agctgcccaa acaacttgat cgtggaagat 60
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<213> Artificial Sequence

27

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<223> Description of Artificial Sequence: Primer
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<223> Description of Artificial Sequence: Synthetic
      peptide
Asp Asn Pro Ala Ser Thr Thr Asn Lys Asp Lys
<210> 10
<211> 9
<212> PRT
<213> Artificial Sequence
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<223> Description of Artificial Sequence: Synthetic
      peptide
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Pro Gln Ala Ser Gly Val Tyr Met Gly
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<211> 16
<212> PRT
<213> Artificial Sequence
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      peptide
Glu Arg Pro Gln Ala Ser Gly Val Tyr Met Gly Asn Leu Thr Ala Gln
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	12 gata acceggegte gaccactaac aaggataage tgttegeagt geeteaggee gtgt atatgggt	60 78
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<210><211><212><212><213>	32	,
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54

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<211> 61
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      polypeptide PV-PV
Val Leu Gly Leu Arg Leu Met Asp Gly Thr Trp Val Ser Met Gln Thr
Ser Asn Glu Thr Lys Trp Cys Pro Pro Gly Gln Leu Ile Asn Leu His
Asp Phe Arg Ser Asp Glu Ile Glu His Leu Val Val Glu Glu Leu Val
         35
Lys Lys Arg Glu Glu Cys Leu Asp Ala Leu Glu Ser Ile
     50
<210> 19
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<212> PRT
<213> Artificial Sequence
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<223> Description of Artificial Sequence: Chimeric
      polypeptide -PVIII
<400> 19
Ser Pro Pro Gly Gln Leu Ile Asn Leu His Asp Phe Arg Ser Asp Glu
Ile Glu His Leu Val Val Glu Glu Leu Val Lys Lys Arg Glu Glu Cys
             20
                                  25
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<210> 20 <211> 64

<212> PRT

Leu Asp Ala Leu Glu Ser Ile

35

<213> Artificial Sequence

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<223> Description of Artificial Sequence: Chimeric
 polypeptide EBL1-PV

<400> 20

Val Pro Gly Met Arg Leu Met Asp Gly Ser Trp Val Ser Leu Gln Lys
1 5 10 15

Thr Glu Ala Pro Glu Trp Cys Ser Asn Ser Pro Pro Gly Gln Leu Ile 20 25 30

Asn Leu His Asp Phe Arg Ser Asp Glu Ile Glu His Leu Val Val Glu 35 40 45

Glu Leu Val Lys Lys Arg Glu Glu Cys Leu Asp Ala Leu Glu Ser Ile 50 55 60

<210> 21

<211> 61

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Chimeric
 polypeptide Mok-PV

<400> 21

Arg Pro Gly Ile Arg Leu Phe Asp Gly Thr Trp Val Ser Phe Thr Lys
1 5 10 15

Pro Asp Val His Val Trp Cys Thr Pro Asn Gln Leu Ile Asn Leu His 20 25 30

Asp Phe Arg Ser Asp Glu Ile Glu His Leu Val Val Glu Glu Leu Val
35 40 45

Lys Lys Arg Glu Glu Cys Leu Asp Ala Leu Glu Ser Ile 50 55 60

<210> 22

<211> 61

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Chimeric
 polypeptide PV-Mok

<400> 22

Val Leu Gly Leu Arg Leu Met Asp Gly Thr Trp Val Ser Met Gln Thr
1 5 10 15

Ser Asn Glu Thr Lys Trp Cys Pro Pro Gly Gln Leu Ile Asn Ile His 20 25 30 Asn Asp Arg Leu Asp Glu Ile Glu His Leu Ile Val Glu Asp Ile Ile 35 40 45

Lys Lys Arg Glu Glu Cys Leu Asp Thr Leu Glu Thr Ile 50 55 60

<210> 23

<211> 61

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Chimeric
 polypeptide Mok-Sad

<400> 23

Lys Pro Gly Ile Arg Leu Phe Asp Gly Thr Trp Val Ser Phe Thr Lys

1 5 10 15

Pro Asp Glu Thr Lys Trp Cys Pro Pro Asp Lys Leu Val Asn Leu His
20 25 30

Asp Phe Arg Ser Asp Glu Ile Glu His Leu Val Val Glu Glu Leu Val
35 40 45

Arg Lys Arg Glu Glu Cys Leu Asp Ala Leu Glu Ser Ile 50 55 60

<210> 24

<211> 524

<212> PRT

<213> Lyssavirus sp.

<220>

<223> PV

<400> 24

Met Val Pro Gln Ala Leu Leu Phe Val Pro Leu Leu Val Phe Pro Leu

1 5 10 15

Cys Phe Gly Lys Phe Pro Ile Tyr Thr Ile Pro Asp Lys Leu Gly Pro
20 25 30

Trp Ser Pro Ile Asp Ile His His Leu Ser Cys Pro Asn Asn Leu Val 35 40 45

Val Glu Asp Glu Gly Cys Thr Asn Leu Ser Gly Phe Ser Tyr Met Glu
50 55 60

Leu Lys Val Gly Tyr Ile Ser Ala Ile Lys Met Asn Gly Phe Thr Cys
65 70 75 80

Thr Gly Val Val Thr Glu Ala Glu Thr Tyr Thr Asn Phe Val Gly Tyr
85 90 95

Val Thr Thr Thr Phe Lys Arg Lys His Phe Arg Pro Thr Pro Asp Ala 100 105 Cys Arg Ala Ala Tyr Asn Trp Lys Met Ala Gly Asp Pro Arg Tyr Glu Glu Ser Leu His Asn Pro Tyr Pro Asp Tyr His Trp Leu Arg Thr Val 130 135 Lys Thr Thr Lys Glu Ser Leu Val Ile Ile Ser Pro Ser Val Ala Asp 150 155 Leu Asp Pro Tyr Asp Arg Ser Leu His Ser Arg Val Phe Pro Gly Gly 165 170 Asn Cys Ser Gly Val Ala Val Ser Ser Thr Tyr Cys Ser Thr Asn His 180 Asp Tyr Thr Ile Trp Met Pro Glu Asn Pro Arg Leu Gly Met Ser Cys 200 Asp Ile Phe Thr Asn Ser Arg Gly Lys Arg Ala Ser Lys Gly Ser Glu 215 Thr Cys Gly Phe Val Asp Glu Arg Gly Leu Tyr Lys Ser Leu Lys Gly 235 Ala Cys Lys Leu Lys Leu Cys Gly Val Leu Gly Leu Arg Leu Met Asp Gly Thr Trp Val Ala Met Gln Thr Ser Asn Glu Thr Lys Trp Cys Pro 260 Pro Gly Gln Leu Val Asn Leu His Asp Phe Arg Ser Asp Glu Ile Glu 280 His Leu Val Val Glu Glu Leu Val Lys Lys Arg Glu Glu Cys Leu Asp 295 Ala Leu Glu Ser Ile Met Thr Thr Lys Ser Val Ser Phe Arg Arg Leu 305 310 315 Ser His Leu Arg Lys Leu Val Pro Gly Phe Gly Lys Ala Tyr Thr Ile Phe Asn Lys Thr Leu Met Glu Ala Asp Ala His Tyr Lys Ser Val Arg 340 345 350 Thr Trp Asn Glu Ile Ile Pro Ser Lys Gly Cys Leu Arg Val Gly Gly Arg Cys His Pro His Val Asn Gly Val Phe Phe Asn Gly Ile Ile Leu 375 Gly Pro Asp Gly Asn Val Leu Ile Pro Glu Met Gln Ser Ser Leu Leu 385

390

395

400

Gln Gln His Met Glu Leu Leu Val Ser Ser Val Ile Pro Leu Met His
405 410 415

Pro Leu Ala Asp Pro Ser Thr Val Phe Lys Asn Gly Asp Glu Ala Glu 420 425 430

Asp Phe Val Glu Val His Leu Pro Asp Val His Glu Arg Ile Ser Gly
435 440 445

Val Asp Leu Gly Leu Pro Asn Trp Gly Lys Tyr Val Leu Leu Ser Ala 450 455 460

Gly Ala Leu Thr Ala Leu Met Leu Ile Ile Phe Leu Met Thr Cys Trp 465 470 475 480

Arg Arg Val Asn Arg Ser Glu Pro Thr Gln His Asn Leu Arg Gly Thr
485
490
495

Gly Arg Glu Val Ser Val Thr Pro Gln Ser Gly Lys Ile Ile Ser Ser 500 505 510

Trp Glu Ser Tyr Lys Ser Gly Gly Glu Thr Gly Leu
515 520

<210> 25

<211> 524

<212> PRT

<213> Lyssavirus sp.

<220>

<223> USA7-BT

<400> 25

Met Ile Pro Gln Ala Leu Leu Phe Val Pro Leu Leu Ile Pro Ser Leu
1 5 10 15

Cys Leu Gly Glu Phe Pro Ile Tyr Thr Ile Pro Asp Lys Leu Gly Pro 20 25 30

Trp Thr Pro Ile Asp Ile His His Leu Ser Cys Pro Asn Asn Leu Val
35 40 45

Ala Glu Asn Asp Gly Cys Thr Ser Leu Ser Gly Phe Ser Tyr Met Glu 50 55 60

Leu Lys Val Gly Tyr Ile Ser Ala Ile Lys Val Asn Gly Phe Thr Cys
65 70 75 80

Thr Gly Val Val Thr Glu Ala Glu Thr Tyr Thr Asn Phe Val Gly Tyr 85 90 95

Val Thr Thr Phe Lys Arg Lys His Phe Arg Pro Met Pro Asp Ala 100 105 110

Cys Arg Ala Ala His Asp Trp Lys Met Ala Gly Asp Pro Arg Tyr Glu 115 120 125

Asp Ser Leu Gln Asn Pro Tyr Pro Asp Tyr His Trp Leu Arg Thr Val 130 135 140 Lys Thr Thr Lys Glu Ser Leu Val Ile Ile Ser Pro Ser Val Ala Asp 150 155 Leu Asp Pro Tyr Asp Lys Ser Leu His Ser Arg Val Phe Pro Ser Gly 170 Lys Cys Leu Gly Ile Thr Val Ser Ser Thr Tyr Cys Ser Thr Asn His 180 Asp Tyr Thr Ile Trp Met Pro Val Glu Pro Arg Leu Gly Thr Ser Cys 200 Asp Ile Phe Thr Asn Ser Arg Gly Lys Arg Ala Ser Lys Gly Gly Arg 210 Val Cys Gly Phe Val Asp Glu Arg Gly Leu Tyr Lys Ser Leu Lys Gly 230 225 Ala Cys Lys Leu Lys Leu Cys Gly Val Pro Gly Ile Arg Leu Met Asp 250 Gly Thr Trp Val Ser Ile Gln Thr Ser Glu Asp Ile Lys Trp Cys Pro 260 Pro Asp Arg Leu Val Asn Leu His Asp Phe His Ser Asp Glu Leu Glu His Leu Val Val Glu Glu Leu Ile Lys Arg Arg Glu Asn Cys Leu Asp 295 Ala Leu Glu Ser Ile Met Thr Thr Lys Ser Val Ser Phe Arg Arg Leu 305 310 Ser His Leu Arg Arg Leu Val Pro Gly Phe Gly Lys Ala Tyr Thr Ile 330 Phe Asn Lys Thr Leu Ile Glu Ala Asp Ala His Tyr Lys Ser Ile Lys 340 345 350 Thr Trp Asn Glu Val Ile Pro Ser Lys Gly Cys Leu Glu Val Gly Gly Lys Cys His Pro Pro Val Asn Gly Val Phe Phe Asn Gly Ile Ile Leu 375 Gly Pro Asp Gly Asn Val Leu Ile Pro Glu Met Gln Ser Ser Leu Leu 385 390 Gln Gln His Met Glu Leu Leu Glu Ser Ser Val Ile Pro Leu Met His 410 Pro Leu Ala Asp Pro Ser Thr Val Phe Lys Glu Gly Asp Glu Ala Glu 430

Asp Phe Val Glu Val His Leu Pro Asp Val His Lys Arg Ile Ser Gly 435 440 445

Val Asp Leu Gly Leu Pro Ser Trp Gly Lys Tyr Leu Leu Met Ser Ala 450 455 460

Gly Ala Leu Ala Ile Leu Ile Leu Ala Ile Val Leu Ile Ile Cys Cys 465 470 475 480

Arg Arg Val Asn Lys Thr Gly Ser Thr Gln Arg Gly His Arg Glu Ser 485 490 495

Arg Gly Lys Met Ser Val Ala Pro Gln Asn Gly Lys Ile Ile Ser Ser 500 505 510

Trp Glu Leu Tyr Lys Arg Glu Ser Glu Thr Gly Leu
515 520

<210> 26

<211> 525

<212> PRT

<213> Lyssavirus sp.

<220>

<223> PL

<400> 26

Met Leu Leu Gln Ile Val Leu Leu Met Ser Leu Met Val Phe Ser Pro 1 5 10 15

Cys Pro Gly Lys Phe Pro Leu Tyr Thr Ile Pro Asp Lys Leu Gly Pro

Trp Ser Pro Ile Asp Ile His His Leu Ser Cys Pro Asn Asn Leu Ile

Val Glu Asp Glu Gly Cys Thr Ser Leu Ser Gly Phe Ser Tyr Met Glu
50 55 60

Leu Lys Val Gly Phe Ile Thr Thr Ile Lys Val Ser Gly Phe Thr Cys
65 70 75 80

Thr Gly Val Val Thr Glu Ser Glu Thr Tyr Thr Asn Phe Phe Gly Tyr 85 90 95

Val Thr Thr Thr Phe Lys Arg Lys His Phe Arg Pro Thr Pro Glu Phe

Cys Arg Asn Ala Tyr Asn Trp Lys Val Ala Gly Asp Pro Arg Tyr Glu

Glu Ser Leu His Asn Pro Tyr Pro Asp Tyr His Trp Leu Arg Thr Val 130 135 140

Thr Thr Thr Lys Glu Ser Leu Leu Ile Ile Ser Pro Ser Val Val Asp 145 150 155 160 Met Asp Pro Tyr Asp Lys Ser Leu His Ser Lys Met Phe Pro Lys Gly 165 170 175

Thr Cys Ser Gly Ala Ser Val Pro Ser Ile Phe Cys Ser Thr Asn His 180 185 190

Asp Tyr Thr Leu Trp Met Pro Glu Asn Pro Lys Pro Gly Met Ser Cys
195 200 205

Asp Ile Phe Thr Thr Ser Lys Gly Lys Lys Ala Ser Lys Gly Gly Lys 210 215 220

Val Cys Gly Phe Val Asp Glu Arg Gly Leu Tyr Lys Ser Leu Lys Gly
225 230 235 240

Ala Cys Lys Leu Lys Leu Cys Gly Ile Ser Gly Leu Arg Leu Met Asp 245 250 255

Gly Ser Trp Val Ser Ile Gln Asn His Glu Glu Ala Lys Trp Cys Ser 260 265 270

Pro Asp Gln Leu Val Asn Ile His Asp Phe His Ser Asp Glu Ile Glu 275 280 285

His Leu Ile Val Glu Glu Leu Val Arg Lys Arg Glu Glu Cys Leu Asp 290 295 300

Ala Leu Glu Ser Ile Met Thr Thr Lys Ser Val Ser Phe Arg Arg Leu 305 310 315 320

Ser His Leu Arg Lys Leu Val Pro Gly Phe Gly Lys Ala Tyr Thr Ile 325 330 335

Val Asn Lys Thr Leu Met Glu Ala Asp Ala His Tyr Asn Gln Val Arg 340 345 350

Thr Trp Asn Glu Ile Ile Pro Ser Lys Gly Cys Leu Lys Val Arg Glu
355 360 365

Arg Cys His Pro Pro Tyr Asn Gly Val Phe Phe Asn Gly Ile Ile Leu 370 375 380

Ser Pro Asp Gly His Val Leu Ile Pro Glu Met Gln Ser Ser Leu Leu 385 390 395 400

Gln Gln His Ile Glu Leu Leu Glu Ser Ser Val Ile Pro Leu Ile His
405 410 415

Pro Leu Ala Asp Pro Ser Thr Val Phe Tyr Arg Asp Asp Glu Ala Glu
420 425 430

Asp Phe Ile Glu Val His Leu Pro Asp Val Gln Lys Gln Val Ser Gly
435
440

Ile Asp Leu Gly Leu Ser Glu Trp Glu Arg Tyr Leu Ile Ile Gly Ala 450 455 460

Ser Ala Val Ile Leu Phe Ala Leu Ala Ile Ile Phe Ala Val Cys Cys 465 470 475 480

Arg Arg Cys Lys Arg Arg Lys Lys Ala Arg Thr Asp Arg Ile Glu Leu
485 490 495

Asp Arg Lys Val Ser Val Thr Ser Gln Ser Gly Lys Val Ile Pro Ser 500 505 510

Trp Glu Ser Tyr Lys Leu Pro Lys Ser His Phe Arg Ser 515 520 525

<210> 27

<211> 524

<212> PRT

<213> Lyssavirus sp.

<220>

<223> EBL1POL

<400> 27

Met Leu Leu Ser Thr Ala Ile Phe Ala Phe Phe Leu Asn Cys Ala Pro 1 5 10 15

Cys Leu Ala Lys Phe Pro Ile Tyr Thr Ile Pro Asp Lys Ile Gly Pro 20 25 30

Trp Ser Pro Ile Asp Ile Asn His Leu Ser Cys Pro Asn Asn Leu Ile 35 40 45

Val Glu Asp Glu Gly Cys Thr Thr Leu Thr Pro Phe Ser Tyr Met Glu 50 55 60

Leu Lys Val Gly Tyr Ile Thr Thr Ile Ile Glu Ser Gly Phe Thr Cys
65 70 75 80

Thr Gly Val Ile Thr Glu Ala Glu Thr Tyr Thr Asn Phe Val Gly Tyr
85 90 95

Val Thr Thr Phe Lys Arg Lys His Phe Arg Pro Thr Val Ser Ala 100 105 110

Cys Arg Asp Ala Tyr Asn Trp Lys Ile Thr Gly Asp Pro Arg Tyr Glu 115 120 125

Glu Ser Leu His Asn Pro Tyr Pro Asp Ser His Trp Leu Arg Thr Val 130 135 140

Lys Thr Thr Lys Glu Ser Leu Leu Ile Ile Ser Pro Ser Val Val Asp 145 150 155 160

Met Asp Ala Tyr Asp Lys Asn Leu Tyr Ser Lys Met Phe Pro Asn Gly 165 170 175

Lys Cys Leu Ala Ser Pro Pro Ser Ala Ile Cys Cys Pro Thr Asn His 180 185 190 Asp Tyr Thr Ile Trp Ile Pro Glu Asn Pro Lys Pro Gly Leu Ser Cys 200 Asp Ile Phe Thr Thr Ser Lys Gly Lys Lys Ala Thr Lys Asp Gly Arg Leu Cys Gly Phe Val Asp Glu Arg Gly Leu Tyr Lys Ser Leu Lys Gly 225 235 Ala Cys Lys Leu Arg Leu Cys Gly Val Pro Gly Met Arg Leu Met Asp Gly Ser Trp Val Ser Leu Gln Lys Thr Glu Ala Pro Glu Trp Cys Ser 265 Pro Asp Gln Leu Val Asn Val His Asp Phe His Thr Asp Glu Ile Glu 275 280 His Leu Val Val Glu Glu Leu Val Lys Lys Arg Glu Glu Cys Leu Asp 295 Ala Leu Glu Thr Ile Ile Thr Thr Lys Ser Ile Ser Phe Arg Arg Leu .315 Ser His Phe Arg Lys Leu Val Pro Gly Phe Gly Lys Ala Tyr Thr Leu Ile Asn Lys Thr Leu Met Glu Ala Asp Ala His Tyr Lys Ser Val Arg Glu Trp Lys Glu Val Ile Pro Ser Lys Gly Cys Leu Met Ala Gly Gly 355 360 Arg Cys His Pro His Tyr Ser Gly Ile Phe Phe Asn Gly Ile Ile Leu Ser Pro Gly Gly Asp Val Leu Ile Pro Glu Met Gln Ser Ala Leu Leu 390 395 Gln Gln His Ile Glu Leu Leu Glu Ser Ser Met Ile Pro Leu Arg His 405 415 Pro Leu Ala Asp Pro Ser Thr Val Phe Arg Lys Asp Asp Glu Ala Glu 425 Asp Phe Val Glu Val His Leu Pro Asp Thr Gln Lys Leu Ile Ser Gly 440 Ile Asp Leu Gly Phe Pro Glu Trp Lys Arg Tyr Phe Leu Ile Gly Ile 450 455 Ser Val Leu Ala Leu Leu Ala Leu Ala Ile Ile Thr Ala Ala Cys Cys 475 Lys Arg Phe Lys Arg Arg Arg Pro Lys Pro Asn Pro Ile Glu Leu

Ile Arg Lys Val Ser Val Thr Ser Gln Ser Gly Arg Ala Ile Pro Ser 500 505 510

Trp Glu Ser Tyr Lys Val Gly Pro Pro Gly Glu Ser 515 520

<210> 28

<211> 524

<212> PRT

<213> Lyssavirus sp.

<220>

<223> EBL1FRA

<400> 28

Met Leu Leu Ser Thr Ala Ile Phe Ala Phe Phe Leu Asn Cys Ala Pro 1 5 10 15

Cys Leu Gly Lys Phe Pro Ile Tyr Thr Ile Pro Asp Lys Ile Gly Pro 20 25 30

Trp Ser Pro Ile Asp Ile Asn His Leu Ser Cys Pro Asn Asn Leu Ile 35 40 45

Val Glu Asp Glu Gly Cys Thr Thr Leu Thr Pro Phe Ser Tyr Met Glu 50 55 60

Leu Lys Val Gly Tyr Ile Thr Thr Ile Lys Ile Glu Gly Phe Thr Cys 65 70 75 80

Thr Gly Val Ile Thr Glu Ala Glu Thr Tyr Thr Asn Phe Val Gly Tyr
85 90 95

Val Thr Thr Thr Phe Lys Arg Lys His Phe Arg Pro Thr Val Ser Ala 100 105 110

Cys Arg Asp Ala Tyr Asn Trp Lys Ile Thr Gly Asp Pro Arg Tyr Glu 115 120 125

Glu Ser Leu His Asn Pro Tyr Pro Asp Ser His Trp Leu Arg Thr Val 130 135 140

Lys Thr Thr Lys Glu Ser Leu Leu Ile Ile Ser Pro Ser Val Val Asp 145 150 155 160

Met Asp Ala Tyr Asp Lys Asn Leu Tyr Ser Lys Met Phe Pro Asn Gly
165 170 175

Lys Cys Leu Ala Ser Pro Pro Ser Ala Thr Cys Cys Pro Thr Asn His 180 185 190

Asp Tyr Thr Ile Trp Ile Pro Glu Asn Pro Lys Pro Gly Leu Ser Cys
195 200 205

Asp Ile Phe Thr Thr Ser Lys Gly Lys Lys Ala Thr Lys Asp Gly Lys 210 215 220

- Leu Cys Gly Phe Val Asp Glu Arg Gly Leu Tyr Lys Ser Leu Lys Gly 225 230 235 240
- Ala Cys Lys Leu Arg Leu Cys Gly Val Pro Gly Met Arg Leu Met Asp 245 250 255
- Gly Ser Trp Val Ser Leu Gln Lys Thr Glu Ala Pro Glu Trp Cys Ser 260 265 270
- Pro Asp Arg Leu Val Asn Ile His Asp Phe His Thr Asp Glu Ile Glu 275 280 285
- His Leu Val Val Glu Glu Leu Val Lys Lys Arg Glu Glu Cys Leu Asp 290 295 300
- Ala Leu Glu Thr Ile Ile Thr Thr Lys Ser Ile Ser Phe Arg Arg Leu 305 310 315 320
- Ser His Phe Arg Lys Leu Val Pro Gly Phe Gly Lys Ala Tyr Thr Leu 325 330 335
- Ile Asn Lys Thr Leu Met Glu Ala Asp Ala His Tyr Lys Ser Val Arg
- Glu Trp Thr Glu Val Ile Pro Ser Lys Gly Cys Leu Met Ala Gly Gly
 355 360 365
- Arg Cys His Pro His Tyr Ser Gly Ile Phe Phe Asn Gly Ile Ile Leu 370 375 380
- Ser Pro Gly Gly Asp Val Leu Ile Pro Glu Met Gln Ser Ala Leu Leu 385 390 395 400
- Gln Gln His Ile Glu Leu Leu Glu Ser Ser Met Ile Pro Leu Arg His 405 410 415
- Pro Leu Ala Asp Pro Ser Thr Val Phe Lys Arg Asp Asp Glu Ala Glu 420 425 430
- Asp Phe Val Glu Val His Leu Pro Asp Thr Gln Lys Leu Ile Ser Gly
 435 440 445
- Ile Asp Leu Gly Phe Pro Glu Trp Lys Arg Tyr Phe Leu Ile Gly Ile
 450 455 460
- Ser Val Leu Ala Leu Leu Ala Leu Ala Ile Ile Thr Ala Ala Cys Cys 465 470 475 480
- Lys Arg Phe Lys Arg Arg Arg Pro Lys Pro Asn Pro Ile Glu Leu 485 490 495
- Ile Arg Lys Val Ser Val Thr Ser Gln Ser Gly Arg Ala Ile Pro Ser 500 505 510
- Trp Glu Ser Tyr Lys Val Gly Thr Thr Ser Glu Ser 515 520

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<210> 29
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<211> 524

<212> PRT

<213> Lyssavirus sp.

<220>

<223> EBL2FIN

<400> 29

Met Pro Phe Gln Thr Val Leu Ser Ala Leu Leu Ser Ala Leu Thr Leu 1 5 10 15

Cys Ala Gly Lys Phe Pro Ile Tyr Thr Ile Pro Asp Lys Leu Gly Pro 20 25 30

Trp Ser Pro Ile Asp Ile His His Leu Ser Cys Pro Thr Asn Met Val 35 40 45

Val Glu Asp Glu Gly Cys Thr Thr Leu Thr Val Phe Ser Tyr Met Glu 50 55 60

Leu Arg Val Gly Tyr Ile Thr Thr Ile Lys Val Asp Gly Phe Thr Cys 65 70 75 80

Thr Gly Val Val Thr Glu Ala Glu Thr Tyr Thr Asn Phe Val Gly Tyr
85 90 95

Val Thr Thr Phe Lys Arg Lys His Phe Arg Pro Ser Pro Ser Ala 100 105 110

Cys Arg Asp Ala Tyr Ser Trp Lys Asn Ala Gly Asp Pro Arg Tyr Glu 115 120 125

Glu Ser Leu His Asn Pro Tyr Pro Asp Ser His Trp Leu Arg Thr Val 130 135 140

Thr Thr Lys Glu Ser Leu Leu Ile Ile Ser Pro Ser Val Val Asn 145 150 155 160

Met Asp Ala Tyr Asp Lys Thr Leu Tyr Ser Lys Ile Phe Leu Asn Gly
165 170 175

Lys Cys Ser Gly Val Ser Gln Val Ser Pro Phe Cys Ser Thr Asn His 180 185 190

Asp Tyr Thr Ile Trp Met Pro Glu Asn Pro Asn Pro Gly Val Ser Cys
195 200 205

Asp Ile Phe Thr Thr Ser Lys Gly Lys Lys Ala Thr Lys Asp Gly Lys 210 215 220

Leu Cys Gly Phe Val Asp Glu Arg Gly Leu Tyr Lys Ser Leu Lys Gly
225 230 235 240

Ala Cys Lys Leu Lys Leu Cys Gly Ile Ser Gly Met Arg Leu Met Asp 245 250 255 Gly Ser Trp Val Ser Ile Gln Asn His Asp Glu Ala Lys Trp Cys Ser 260 265 270

Pro Asp Gln Leu Val Asn Ile His Asp Phe His Ser Asp Glu Val Glu 275 280 285

His Leu Ile Ala Glu Glu Leu Val Lys Lys Arg Glu Glu Cys Leu Asp 290 295 300

Ala Leu Glu Ser Ile Met Thr Thr Lys Ser Ile Ser Phe Arg Arg Leu 305 310 315 320

Ser His Leu Arg Lys Leu Val Pro Gly Phe Gly Lys Ala Tyr Thr Ile 325 330 335

Ile Asn Lys Thr Leu Met Glu Ala Asp Ala His Tyr Lys Ser Ile Arg
340 345 350

Glu Trp Thr Asp Val Ile Pro Ser Lys Gly Cys Leu Met Ala Gly Gly
355 360 365

Arg Cys Tyr Pro His His Asn Gly Val Phe Phe Asn Gly Ile Ile Leu 370 375 380

Ser Pro Asp Gly His Val Leu Ile Pro Glu Met Gln Ser Ala Met Leu 385 390 395 400

Gln Gln His Ile Glu Leu Leu Glu Ser Ser Val Ile Pro Leu Met His
405 410 415

Pro Leu Ala Asp Pro Ser Thr Ile Phe Lys Lys Asp Asp Gly Ala Glu
420 425 430

Asp Phe Val Glu Val His Leu Pro Asp Val Gln Lys Gln Ile Ser Gly
435 440 445

Ile Asp Leu Gly Leu Pro Glu Trp Lys Arg Tyr Phe Leu Ile Gly Val 450 455 460

Ser Ala Leu Ala Leu Leu Ala Leu Met Ile Phe Ile Ala Ala Cys Cys 465 470 475 480

Lys Arg Val Lys His Lys Lys Arg Ala Lys Pro Asn Pro Val Glu Leu 485 490 495

Ile Arg Lys Val Ser Val Thr Ser Gln Ser Gly Arg Pro Ile Pro Ser 500 505 510

Trp Glu Ser Tyr Lys Val Glu Thr Gly Gly Gln Ser 515 520

<210> 30

<211> 524

<212> PRT

<213> Lyssavirus sp.

<220>

<223> EBL2HOL

<400> 30

Met Pro Phe Gln Ala Val Leu Ser Ala Leu Leu Ser Ala Leu Thr Leu 1 5 10 15

Cys Val Gly Lys Phe Pro Ile Tyr Thr Ile Pro Asp Lys Leu Gly Pro 20 25 30

Trp Ser Pro Ile Asp Ile His His Leu Ser Cys Pro Thr Asn Met Val 35 40 45

Val Glu Asp Glu Gly Cys Thr Thr Leu Thr Val Phe Ser Tyr Met Glu 50 55 60

Leu Lys Val Gly Tyr Ile Thr Thr Ile Lys Val Asn Glu Phe Thr Cys 65 70 75 80

Thr Gly Val Val Thr Glu Ala Glu Thr Tyr Thr Asn Phe Val Gly Tyr
85 90 95

Val Thr Thr Phe Lys Arg Lys Asp Phe Arg Pro Ser Pro Ser Ala 100 105 110

Cys Arg Asp Ala Tyr Ser Cys Lys Thr Ala Gly Asp Pro Arg Tyr Glu 115 120 125

Glu Ser Leu His Asn Pro Tyr Pro Asp Ser His Trp Leu Thr Cys Thr 130 135 140

Thr Thr Thr Lys Glu Ser Val Leu Ile Ile Ser Pro Ser Val Ala Asp 145 150 155 160

Met Asp Ala Tyr Asp Lys Thr Leu Tyr Ser Lys Ile Phe Leu Asn Gly
165 170 175

Lys Cys Ser Gly Val Ser Gln Val Ser Pro Phe Cys Ser Thr Asn His 180 185 190

Asp Tyr Thr Ile Trp Met Pro Glu Asn Pro Asn Pro Gly Val Ser Cys
195 200 205

Asp Ile Phe Thr Thr Ser Lys Gly Lys Lys Ala Thr Lys Asp Gly Lys 210 215 220

Leu Cys Gly Phe Val Asp Glu Arg Gly Leu Tyr Lys Ser Leu Lys Gly 225 230 235 240

Ala Cys Lys Leu Lys Leu Cys Gly Ile Ser Gly Met Arg Leu Met Asp 245 250 255

Gly Ser Trp Val Ser Ile Gln Asn His Asp Glu Ala Lys Trp Cys Ser 260 265 270

Pro Asp Gln Leu Val Asn Ile His Asp Phe His Ser Asp Glu Val Glu 275 280 285

His Leu Ile Ala Glu Glu Leu Val Lys Lys Arg Glu Glu Cys Leu Asp 290 295 300

Ala Leu Glu Ser Ile Met Thr Thr Lys Ser Ile Ser Phe Arg Arg Leu 305 310 315 320

Ser His Leu Arg Lys Leu Val Pro Gly Phe Gly Lys Ala Tyr Thr Val 325 330 335

Ile Asn Lys Thr Leu Met Glu Ala Asp Ala His Tyr Lys Ser Ile Arg 340 345 350

Glu Trp Thr Asp Val Ile Pro Ser Lys Gly Cys Leu Met Ala Gly Gly 355 360 365

Arg Cys Tyr Pro His His Asn Gly Val Phe Phe Asn Gly Ile Ile Leu 370 375 380

Ser Pro Asp Gly His Val Leu Ile Pro Glu Met Gln Ser Ala Met Leu 385 390 395 400

Gln Gln His Ile Glu Leu Leu Glu Ser Ser Val Ile Pro Leu Met His 405 410 415

Pro Leu Ala Asp Pro Ser Thr Ile Phe Lys Lys Asp Asp Gly Ala Glu
420 425 430

Asp Phe Val Glu Val His Leu Pro Asp Val Gln Lys Gln Ile Ser Gly
435 440 445

Ile Asp Leu Gly Leu Pro Glu Trp Lys Arg Tyr Phe Leu Ile Gly Val 450 455 460

Ser Ala Leu Ala Phe Leu Ala Leu Met Ile Phe Ile Ala Ala Cys Cys 465 470 475 480

Arg Arg Val Lys Arg Lys Lys Arg Ala Lys Pro Asn Pro Val Glu Leu 485 490 495

Ile Arg Lys Val Ser Val Thr Ser Gln Ser Gly Arg Pro Ile Pro Ser 500 505 510

Trp Glu Ser Tyr Lys Val Glu Thr Gly Gly Gln Ser 515 520

<210> 31

<211> 533

<212> PRT

<213> Lyssavirus sp.

<220>

<223> Duv1SAF

<400> 31

Met Pro Leu Asn Ala Val Ile Phe Thr Leu Leu Leu Arg Cys Ser Ile
1 5 10 15

- Cys Leu Gly Lys Phe Pro Phe Tyr Thr Ile Pro Asp Lys Leu Gly Pro
 20 25 30
- Trp Ser Pro Ile Asp Ile His His Leu Ser Cys Pro Asn Asn Leu Val 35 40 45
- Val Glu Asp Glu Gly Cys Thr Thr Leu Thr Pro Phe Ser Tyr Met Glu
 50 55 60
- Leu Lys Val Gly Tyr Ile Thr Ser Ile Lys Val Ser Gly Phe Thr Cys
 65 70 75 80
- Thr Gly Val Val Thr Glu Ala Glu Thr Tyr Thr Asn Phe Val Gly Tyr
 85 90 95
- Val Thr Thr Thr Phe Arg Arg Arg His Phe Arg Pro Ser Val Asn Ser 100 105 110
- Cys Arg Asp Ala Tyr Asn Trp Lys Ile Ala Gly Asp Pro Arg Tyr Glu 115 120 125
- Glu Ser Leu His Asn Pro Tyr Pro Asp Ser His Trp Leu Arg Thr Val 130 135 140
- Lys Thr Thr Lys Glu Ser Leu Leu Ile Ile Ser Pro Ser Val Ala Asp 145 150 155 160
- Met Asp Ala Tyr Asp Lys Lys Leu Tyr Ser Lys Met Phe Pro Asn Gly
 165 170 175
- Arg Cys Ser Glu Ile Ser Pro Gly Ser Pro Phe Cys Pro Thr Asn His 180 185 190
- Glu Tyr Thr Ile Trp Met Pro Glu Ser Ser Asn Pro Gly Ile Ser Cys 195 200 205
- Asp Ile Phe Thr Arg Ser Met Gly Lys Lys Ala Thr Lys Asp Gly Gln 210 215 220
- Leu Cys Gly Phe Val Asp Glu Arg Gly Leu Tyr Lys Ser Leu Lys Gly 225 230 235 240
- Ala Cys Arg Leu Arg Leu Cys Gly Ile Ser Gly Leu Arg Leu Met Asp 245 250 255
- Gly Ser Trp Val Ser Leu Pro Gln Val Asn Asn Ser Glu Trp Cys Ser 260 265 270
- Pro Asp Gln Leu Val Asn Ile His Asp Phe His Ser Asp Glu Ile Glu 275 280 285
- His Leu Val Ala Asp Glu Leu Val Lys Lys Arg Glu Asp Cys Leu Asp 290 295 300
- Ala Leu Glu Thr Ile Ile Phe Thr Lys Ser Ile Ser Phe Arg Arg Leu 305 310 315 320

Ser Arg Leu Arg Lys Leu Val Pro Gly Phe Gly Lys Ala Tyr Thr Ile 325 330 335

Ile Asn Arg Thr Leu Met Glu Ala Glu Ala His Tyr Lys Ser Val Arg 340 345 350

Glu Trp Lys Glu Ile Ile Pro Ser Lys Gly Cys Leu Lys Ala Gly Gly 355 360 365

Arg Cys Tyr Pro His His Asn Ile Val Phe Phe Asn Gly Ile Ile Leu 370 375 380

Gly Pro Gly Gly Lys Ile Leu Ile Pro Glu Met Gln Ser Ala Leu Leu 385 390 395 400

Gln Gln His Ile Glu Leu Leu Glu Ser Ser Val Val Pro Leu Lys His
405 410 415

Pro Leu Ala Asp Pro Ser Thr Val Phe Lys Asn Asp Asp Glu Ala Glu
420 425 430

Ser Phe Val Asp Val His Leu Pro Asp Thr Asn Gln Lys Ile Ser Gly
435 440 445

Ile Asp Leu Gly Leu Pro Glu Trp Lys Arg Tyr Phe Leu Ile Gly Val 450 455 460

Ser Ala Val Ala Leu Leu Ala Leu Ser Ile Ile Met Arg Val Cys Cys 465 470 475 480

Lys Arg Phe Lys Asn Arg Arg Lys Ser Lys Pro Ser Pro Val Glu Leu 485 490 495

Thr Arg Lys Val Ser Val Ile Ser Lys Gly Asn Gly Pro Val Pro Ser 500 505 510

Trp Glu Ser Tyr Lys Glu Gly Thr Thr Gly Asp Val Arg Asn Thr Thr 515 520 525

Pro Ser Thr Arg Glu 530

<210> 32

<211> 533

<212> PRT

<213> Lyssavirus sp.

<220>

<223> Duv2SAF

<400> 32

Met Pro Leu Asn Ala Val Ile Phe Thr Leu Leu Leu Arg Cys Ser Ile 1 5 10 15

Cys Leu Gly Lys Phe Pro Phe Tyr Thr Ile Pro Asp Lys Leu Gly Pro 20 25 30

- Trp Ser Pro Ile Asp Ile His His Leu Ser Cys Pro Asn Asn Leu Val 35 40 45
- Val Glu Asp Glu Gly Cys Thr Thr Leu Thr Pro Phe Ser Tyr Met Glu 50 55 60
- Leu Lys Val Gly Tyr Ile Thr Ser Ile Lys Val Ser Gly Phe Thr Cys
 65. 70 75 80
- Thr Gly Val Val Thr Glu Ala Glu Thr Tyr Thr Asn Phe Val Gly Tyr 85 90 95
- Val Thr Thr Phe Arg Arg Arg His Phe Arg Pro Ser Val Asn Ser 100 105 110
- Cys Arg Asp Ala Tyr Asn Trp Lys Ile Ala Gly Asp Pro Arg Tyr Glu 115 120 125
- Glu Ser Leu His Asn Pro Tyr Pro Asp Ser His Trp Leu Arg Thr Val 130 135 140
- Lys Thr Thr Lys Glu Ser Leu Leu Ile Ile Ser Pro Ser Val Ala Asp 145 150 155 160
- Met Asp Ala Tyr Asp Lys Lys Leu Tyr Ser Lys Met Phe Pro Asn Gly
 165 170 175
- Arg Cys Ser Glu Ile Ser Pro Gly Ser Pro Phe Cys Pro Thr Asn His 180 185 190
- Glu Tyr Thr Ile Trp Met Pro Glu Ser Ser Asn Pro Gly Ile Ser Cys
- Asp Ile Phe Thr Arg Ser Met Gly Lys Lys Ala Thr Lys Asp Gly Gln
- Leu Cys Gly Phe Val Asp Glu Arg Gly Leu Tyr Lys Ser Leu Lys Gly 225 230 235 240
- Ala Cys Arg Leu Arg Leu Cys Gly Ile Ser Gly Leu Arg Leu Met Asp 245 250 255
- Gly Ser Trp Val Ser Leu Pro Gln Val Asn Asn Ser Glu Trp Cys Ser
- Pro Asp Gln Leu Val Asn Ile His Asp Phe His Ser Asp Glu Ile Glu 275 280 285
- His Leu Val Ala Asp Glu Leu Val Lys Lys Arg Glu Asp Cys Leu Asp 290 295 300
- Ala Leu Glu Thr Ile Leu Phe Thr Lys Ser Ile Ser Phe Arg Arg Leu 305 310 315 320
- Ser His Leu Arg Lys Leu Val Pro Gly Phe Gly Lys Ala Tyr Thr Ile 325 330 335

Ile Asn Arg Thr Leu Met Glu Ala Glu Ala His Tyr Lys Ser Val Arg 340 345 350

Glu Trp Lys Glu Ile Ile Pro Ser Lys Gly Cys Leu Lys Ala Gly Gly 355 360 365

Arg Cys Tyr Pro His His Asn Gly Ile Phe Phe Asn Gly Ile Ile Leu 370 375 380

Gly Pro Gly Glu Ile Leu Ile Pro Glu Met Gln Ser Ala Leu Leu 385 390 395 400

Gln Gln His Ile Glu Leu Leu Glu Ser Ser Val Val Pro Leu Lys His
405 410 415

Pro Leu Ala Asp Pro Ser Thr Val Phe Lys Asn Asp Asp Glu Ala Glu 420 425 430

Ser Phe Val Asp Val His Leu Pro Asp Thr Asn Gln Lys Ile Ser Gly 435 440 445

Ile Asp Leu Gly Leu Pro Glu Trp Lys Arg Tyr Phe Leu Ile Gly Val 450 455 460

Ser Ala Val Ala Leu Leu Ala Leu Ser Ile Ile Ile Ala Val Cys Cys 465 470 475 480

Lys Arg Phe Arg Lys Arg Lys Ser Lys Pro Gly Pro Val Glu Leu 485 490 495

Thr Arg Lys Val Ser Val Ile Ser Lys Gly Asn Gly Pro Val Pro Ser 500 505 510

Trp Glu Ser Tyr Lys Glu Gly Thr Thr Gly Asp Val Arg Asn Thr Thr 515 520 525

Pro Ser Thr Arg Glu 530

<210> 33

<211> 522

<212> PRT

<213> Lyssavirus sp.

<220>

<223> Lag1NGA

<400> 33

Met Ser Gln Leu Asn Leu Ile Leu Phe Phe Cys Val Ile Ile Val Leu 1 5 10 15

Ser Val Glu Asp Phe Pro Leu Tyr Thr Ile Pro Glu Lys Ile Gly Pro 20 25 30

Trp Thr Pro Ile Asp Leu Ile His Leu Ser Cys Pro Asn Asn Leu Gln
35 40 45

- Ser Glu Asp Glu Gly Cys Gly Thr Ser Ser Ser Val Ser Tyr Val Glu 50 55 60
- Leu Lys Thr Gly Tyr Leu Thr His Gln Lys Val Ser Gly Phe Thr Cys 65 70 75 80
- Thr Gly Val Val Asn Glu Ala Val Thr Tyr Thr Asn Phe Val Gly Tyr 85 90 95
- Val Thr Thr Thr Phe Lys Arg Lys His Phe Lys Pro Thr Ala Leu Ala 100 105 110
- Cys Arg Asp Ala Tyr His Trp Lys Ile Ser Gly Asp Pro Arg Tyr Glu 115 120 125
- Glu Ser Leu His Thr Pro Tyr Pro Asp Asn Ser Trp Leu Arg Thr Val 130 135 140
- Thr Thr Thr Lys Glu Ser Leu Val Ile Ile Ser Pro Ser Ile Val Glu 145 150 155 160
- Met Asp Val Tyr Ser Arg Thr Leu His Ser Pro Met Phe Pro Thr Gly
 165 170 175
- Thr Cys Ser Arg Phe Tyr Pro Ser Ser Pro Ser Cys Ala Thr Asn His 180 185 190
- Asp Tyr Thr Leu Trp Leu Pro Asp Asp Pro Asn Leu Ser Leu Ala Cys
 195 200 205
- Asp Ile Phe Val Thr Ser Thr Gly Lys Lys Ser Met Asn Gly Ser Arg 210 215 220
- Met Cys Gly Phe Thr Asp Glu Arg Gly Tyr Tyr Arg Thr Ile Lys Gly 225 230 235
- Ala Cys Lys Leu Thr Leu Cys Gly Lys Pro Gly Leu Arg Leu Phe Asp 245 250 255
- Gly Thr Trp Ile Ser Phe Thr Arg Pro Glu Val Thr Thr Trp Cys Leu 260 265 270
- Pro Asn Gln Leu Val Asn Ile His Asn Asn Arg Ile Asp Glu Val Glu 275 280 285
- His Leu Ile Val Glu Asp Leu Ile Arg Lys Arg Glu Glu Cys Leu Asp 290 295 300
- Thr Leu Glu Thr Val Leu Met Ser Lys Ser Ile Ser Phe Arg Arg Val 305 310 315 320
- Ser His Phe Arg Lys Leu Val Pro Gly Tyr Gly Lys Ala Tyr Thr Ile 325 330 335
- Leu Asn Gly Ser Leu Ile Gln Thr Asn Val His Tyr Leu Lys Val Asp 340 345 350

Asn Trp Ser Glu Ile Leu Pro Ser Lys Gly Cys Leu Lys Ile Asn Asn 355 360 365

Gln Cys Val Ala His Asp Glu Gly Val Phe Phe Asn Gly Ile Ile Lys 370 375 380

Gly Pro Asp Gly His Ile Leu Ile Pro Glu Met Gln Ser Ser Leu Trp 385 390 395 400

Lys Gln His Met Asp Leu Phe Lys Ala Ala Val Phe Pro Leu Arg His 405 410 415

Pro Leu Ile Glu Pro Gly Ser Leu Phe Asn Lys Asp Gly Asp Ala Asp 420 425 430

Glu Phe Val Asp Val His Met Pro Asp Val His Lys Leu Val Ser Asp 435 440 445

Val Asp Leu Gly Leu Pro Asp Trp Ser Leu Tyr Ala Leu Ile Gly Ala 450 455 460

Thr Ile Ile Ala Phe Phe Ile Leu Ile Cys Leu Ile Arg Ile Cys Cys 465 470 475 480

Lys Lys Arg Gly Arg Arg Asn Ser Pro Thr Asn Arg Pro Asp Leu Pro 485 490 495

Ile Gly Leu Ser Thr Thr Pro Gln Pro Lys Ser Lys Val Ile Ser Ser 500 505 510

Trp Glu Ser Tyr Lys Gly Thr Ser Asn Val 515 520

<210> 34

<211> 522

<212> PRT

<213> Lyssavirus sp.

<220>

<223> Lag2CAR

<400> 34

Met Ser Gln Leu Ile Leu Ile Pro Phe Leu Cys Val Val Ile Val Ile 1 5 10 15

Ser Val Gly Asp Phe Pro Leu Tyr Thr Ile Pro Glu Lys Ile Gly Thr 20 . 25 30

Trp Thr Pro Ile Asp Leu Ile His Leu Ser Cys Pro Asn Asn Leu Leu 35 40 45

Ser Glu Asp Asp Gly Cys Ser Asn Thr Ala Thr Phe Asn Tyr Ile Glu 50 55 60

Leu Lys Thr Gly Tyr Leu Thr His Gln Lys Val Ser Gly Phe Thr Cys 65 70 75 80

- Thr Gly Val Val Asn Glu Ala Val Thr Tyr Thr Asn Phe Val Gly Tyr 85 90 95
- Val Thr Thr Thr Phe Lys Arg Lys His Phe Lys Pro Thr Ala Leu Ala 100 105 110
- Cys Arg Asp Ala Phe His Trp Lys Ile Ser Gly Asp Pro Arg Tyr Glu 115 120 125
- Glu Ser Leu His Thr Pro Tyr Pro Asp Asn Ser Trp Leu Arg Thr Val 130 135 140
- Thr Thr Thr Lys Glu Ser Leu Leu Ile Ile Ser Pro Ser Ile Val Glu 145 150 155 160
- Met Asp Val Tyr Ser Arg Thr Leu His Ser Pro Met Phe Pro Gly Gly 165 170 175
- Val Cys Ser Lys Phe Tyr Pro Ser Ser Pro Ser Cys Pro Thr Asn His 180 185 190
- Asp Tyr Thr Leu Trp Leu Pro Glu Asp Ala Asn Leu Ser Met Ala Cys 195 200 205
- Asp Ile Phe Ile Thr Ser Thr Gly Lys Lys Ser Met Asn Gly Ser Arg 210 215 220
- Met Cys Gly Phe Thr Asp Glu Arg Gly Phe Tyr Arg Thr Leu Lys Gly
- Ala Cys Lys Leu Thr Leu Cys Gly Lys Pro Gly Leu Arg Leu Tyr Asp 245 250 255
- Gly Thr Trp Val Ser Phe Thr Arg Pro Glu Ile Asn Val Trp Cys Ser 260 265 270
- Pro Asn Gln Leu Val Asn Val His Asn Asn Arg Leu Asp Glu Ile Glu 275 280 285
- His Leu Ile Val Gly Asp Leu Ile Arg Lys Arg Glu Glu Cys Leu Asp 290 295 300
- Thr Leu Glu Thr Ile Leu Met Ser Lys Ser Ile Ser Phe Arg Arg Leu 305 310 315 320
- Ser His Phe Arg Lys Leu Val Pro Gly Tyr Gly Lys Ala Tyr Thr Ile 325 330 335
- Ile Asn Gly Ser Leu Met Glu Thr Asn Val His Tyr Leu Arg Val Asp 340 345 350
- Ser Trp Asn Asp Ile Leu Pro Ser Lys Gly Cys Leu Lys Met Asn Lys 355 360 365
- Gln Cys Val Asp Ser Tyr Arg Gly Val Phe Phe Asn Gly Ile Ile Lys 370 375 380

Gly His Asp Gly His Ile Leu Ile Pro Glu Met Gln Ser Ser Leu Leu 385 390 395 400

Lys Gln His Met Asp Leu Leu Lys Ala Ala Val Phe Pro Leu Arg His
405 410 415

Pro Leu Ile Asp Gln Asn Ser Leu Phe Lys Lys Asp Gly Asp Ala Asp 420 425 430

Asp Phe Val Glu Val His Met Pro Asp Ile Gln Lys Leu Ile Ser Asp 435 440 445

Val Asp Leu Gly Leu Pro Ser Trp Gly Leu Tyr Val Met Ile Gly Ala 450 455 460

Ala Val Ile Ala Phe Leu Val Leu Ile Cys Leu Ile Arg Ile Cys Cys 465 470 475 480

Lys Lys Lys Thr Arg Thr Arg Thr Ser Met Glu Arg Pro Asp Pro Pro 485 490 495

Ile Ser Leu Ser Thr Thr Pro Gln Ser Arg Ala Lys Val Val Ser Ser 500 505 510

Trp Glu Ser Tyr Lys Gly Ser Ser Asn Val 515 520

<210> 35

<211> 523

<212> PRT

<213> Lyssavirus sp.

<220>

<223> Mok3ETP

<400> 35

Met Asn Ile Pro Cys Phe Ala Val Ile Leu Ser Leu Ala Thr Thr His 1 5 10 15

Cys Leu Glu Lys Phe Leu Ile Tyr Thr Ile Pro Glu Lys Ile Glu Lys 20 25 30

Trp Thr Pro Ile Asp Met Ile His Leu Ser Cys Pro Asn Asn Met Leu 35 40 45

Ser Glu Glu Glu Gly Cys Asn Thr Glu Ser Pro Phe Thr Tyr Phe Glu 50 55 60

Leu Lys Ser Gly Tyr Leu Ala His Gln Lys Val Pro Gly Phe Thr Cys 65 70 75 80

Thr Gly Val Val Asn Glu Ala Glu Thr Tyr Thr Asn Phe Val Gly Tyr
85 90 95

Val Thr Thr Thr Phe Lys Arg Lys His Phe Lys Pro Thr Val Ala Ala 100 105 110

- Cys Arg Asp Ala Tyr Asn Trp Lys Val Ser Gly Asp Pro Arg Tyr Glu 115 120 125
- Glu Ser Leu His Thr Pro Tyr Pro Asp Ser Ser Trp Leu Arg Thr Val 130 135 140
- Thr Thr Thr Lys Glu Ala Leu Leu Ile Ile Ser Pro Ile Val Glu Asp 145 150 155 160
- Met Ile Ala Gly Arg Lys Thr Leu His Ser Pro Met Phe Pro Ser Gly
 165 170 175
- Lys Cys Ser Lys Leu Tyr Pro Ser Val Pro Ser Cys Thr Thr Asn His 180 185 190
- Asp Tyr Thr Leu Trp Leu Pro Glu Asp Ser Ser Leu Ser Leu Ile Cys 195 200 205
- Asp Ile Phe Thr Ser Ser Ser Gly Gln Lys Ala Met Asn Gly Ser Arg 210 215 220
- Ile Cys Gly Phe Lys Asp Glu Arg Gly Phe Tyr Arg Ser Leu Lys Gly 225 230 235 240
- Ser Cys Lys Leu Thr Leu Cys Gly Lys Pro Gly Ile Arg Leu Phe Asp 245 250 255
- Gly Thr Trp Val Ser Phe Thr Lys Pro Asp Val His Val Trp Cys Thr 260 265 270
- Pro Asn Gln Leu Val Asn Ile His Asn Asp Arg Leu Asp Glu Val Glu 275 280 285
- His Leu Ile Val Asp Asp Ile Ile Lys Lys Arg Glu Glu Cys Leu Asp 290 295 300
- Thr Leu Glu Thr Ile Leu Met Ser Gln Ser Val Ser Phe Arg Arg Leu 305 310 315 320
- Ser His Phe Arg Lys Leu Val Pro Gly Tyr Gly Lys Ala Tyr Thr Ile 325 330 335
- Leu Asn Gly Ser Leu Met Glu Thr Asn Val Tyr Tyr Lys Arg Val Asp 340 345 350
- Arg Trp Ala Asp Ile Leu Pro Ser Arg Gly Cys Leu Lys Val Gly Gln 355 360 365
- Gln Cys Met Asp Pro Val Lys Asn Leu Val Phe Phe Asn Gly Ile Ile 370 375 380
- Lys Gly Pro Asp Gly Gln Ile Leu Ile Pro Glu Met Gln Ser Glu Gln 385 390 395 400
- Leu Lys Gln His Met Asp Leu Leu Lys Ala Ala Met Phe Pro Leu Arg 405 410 415

His Pro Leu Ile Asn Arg Glu Ala Val Phe Lys Lys Asp Gly Asn Ala 420 425 430

Asp Asp Phe Val Asp Leu His Met Pro Asp Val Gln Lys Ser Val Ser 435 440 445

Asp Val Asp Leu Gly Leu Pro His Trp Gly Phe Trp Leu Leu Val Gly
450 455 460

Ala Thr Val Val Ala Phe Val Val Leu Ala Cys Leu Leu Arg Val Cys 465 470 475 480

Cys Arg Arg Met Arg Arg Arg Ser Leu Arg Ala Thr Gln Asp Ile 485 490 495

Pro Leu Ser Val Ala Pro Ala Pro Val Pro Arg Ala Lys Val Val Ser 500 505 510

Ser Trp Glu Ser Ser Lys Gly Leu Pro Gly Thr 515 520

<210> 36

<211> 523

<212> PRT

<213> Lyssavirus sp.

<220>

<223> Mok2ZIM

<400> 36

Met Asn Ile Pro Cys Phe Val Val Ile Leu Ser Leu Ala Thr Thr His

1 5 10 15

Ser Leu Gly Glu Phe Pro Leu Tyr Thr Ile Pro Glu Lys Ile Glu Lys 20 25 30

Trp Thr Pro Ile Asp Met Ile His Leu Ser Cys Pro Asn Asn Leu Leu 35 40 45

Ser Glu Glu Glu Gly Cys Asn Ala Glu Ser Ser Phe Thr Tyr Phe Glu 50 55 60

Leu Lys Ser Gly Tyr Leu Ala His Gln Lys Val Pro Gly Phe Thr Cys
65 70 75 80

Thr Gly Val Val Asn Glu Ala Glu Thr Tyr Thr Asn Phe Val Gly Tyr 85 90 95

Val Thr Thr Thr Phe Lys Arg Lys His Phe Arg Pro Thr Val Ala Ala 100 105 110

Cys Arg Asp Ala Tyr Asn Trp Lys Val Ser Gly Asp Pro Arg Tyr Glu 115 120 125

Glu Ser Leu His Thr Pro Tyr Pro Asp Ser Ser Trp Leu Arg Thr Val 130 135 140

- Thr Thr Lys Glu Ser Leu Leu Ile Ile Ser Pro Ser Ile Val Glu Met Asp Ile Tyr Gly Arg Thr Leu His Ser Pro Met Phe Pro Ser Gly 170 Val Cys Ser Asn Val Tyr Pro Ser Val Pro Ser Cys Glu Thr Asn His 180 Asp Tyr Thr Leu Trp Leu Pro Glu Asp Pro Ser Leu Ser Leu Val Cys 200 Asp Ile Phe Thr Ser Ser Asn Gly Lys Lys Ala Met Asn Gly Ser Arg 215 Ile Cys Gly Phe Lys Asp Glu Arg Gly Phe Tyr Arg Ser Leu Lys Gly 225 Ala Cys Lys Leu Thr Leu Cys Gly Arg Pro Gly Ile Arg Leu Phe Asp Gly Thr Trp Val Ser Phe Thr Lys Pro Asp Val His Val Lys Thr Cys 265 Asn Pro Asp Ile Leu Val Asn Ile His Asn Asp Arg Leu Asp Glu Ile 275 280 Glu His Leu Ile Val Glu Asp Ile Ile Lys Lys Arg Glu Glu Cys Leu 295 Asp Thr Leu Glu Thr Ile Leu Met Ser Gln Ser Val Ser Phe Arg Arg 305 Leu Ser His Phe Arg Lys Leu Val Pro Gly Tyr Gly Lys Ala Tyr Thr Ile Leu Asn Gly Ser Leu Met Glu Thr Asn Val Tyr Tyr Lys Arg Val 345 Asp Lys Trp Ala Asp Ile Leu Pro Ser Lys Gly Cys Leu Lys Val Gly 355 360 Gln Gln Cys Met Glu Pro Val Lys Gly Val Leu Phe Asn Gly Ile Ile 375 Lys Gly Pro Asp Gly Gln Ile Leu Ile Pro Glu Met Gln Ser Glu Gln 395 385 390
- His Pro Leu Ile Ser Arg Glu Ala Val Phe Lys Lys Asp Gly Asp Ala 420 425 430

Leu Lys Gln His Met Asp Leu Leu Lys Ala Ala Val Phe Pro Leu Arg

405

Asp Asp Phe Val Asp Leu His Met Pro Asp Val His Lys Ser Val Ser 435 440 445 Asp Val Asp Leu Gly Leu Pro His Trp Gly Phe Trp Met Leu Ile Gly 450 455 460

Ala Thr Ile Val Ala Phe Val Val Leu Val Cys Leu Leu Arg Val Cys 465 470 475 480

Cys Lys Arg Val Arg Arg Arg Ser Gly Arg Ala Thr Gln Glu Ile 485 490 495

Pro Leu Ser Phe Pro Ser Ala Pro Val Pro Arg Ala Lys Val Val Ser 500 505 510

Ser Trp Glu Ser Tyr Lys Gly Leu Pro Gly Thr 515 520

<210> 37

<211> 129

<212> PRT

<213> Lyssavirus sp.

<220>

<223> Lag3SAF

<400> 37

Cys Leu Asp Thr Leu Glu Thr Ile Leu Met Ser Lys Ser Ile Ser Phe 1 5 10 15

Arg Arg Leu Ser His Phe Arg Lys Leu Val Pro Gly Tyr Gly Lys Ala

Tyr Thr Ile Ile Asn Gly Ser Leu Met Glu Thr Asn Val His Tyr Leu
35 40 45

Arg Val Asp Ser Trp Asn Asp Ile Leu Pro Ser Lys Gly Cys Leu Lys 50 55 60

Met Asn Lys Gln Cys Val Asp Ser Tyr Arg Gly Val Phe Phe Asn Gly 65 70 75 80

Ile Ile Lys Gly Leu Asp Gly His Ile Leu Ile Pro Glu Met Gln Ser 85 90 95

Ser Leu Leu Lys Gln His Met Asp Leu Leu Lys Ala Ala Val Phe Pro 100 105 110

Leu Arg His Pro Leu Ile Asp Gln Asn Ser Leu Phe Lys Lys Asp Gly
115 120 125

Asp.

<210> 38

<211> 129

<212> PRT

<213> Lyssavirus sp.

<220>

<223> Lag4SAF

<400> 38

Cys Leu Asp Thr Leu Glu Thr Ile Leu Met Ser Lys Ser Ile Ser Phe 1 5 10 15

Arg Arg Leu Ser His Phe Arg Lys Leu Val Pro Gly Tyr Gly Lys Ala 20 25 30

Tyr Thr Ile Ile Asn Gly Ser Leu Met Glu Thr Asn Val His Tyr Leu
35 40 45

Arg Val Asp Ser Trp Asn Asp Ile Leu Pro Ser Lys Gly Cys Leu Lys 50 55 60

Met Asn Lys Gln Cys Val Asp Ser Tyr Arg Gly Val Phe Phe Asn Gly 65 70 75 80

Ile Ile Lys Gly Leu Asp Gly His Ile Leu Ile Pro Glu Met Gln Ser 85 90 95

Ser Leu Leu Lys Gln His Met Asp Leu Leu Lys Ala Ala Val Phe Pro 100 105 110

Leu Arg His Pro Leu Ile Asp Gln Asn Ser Leu Phe Lys Lys Asp Gly
115 120 125

Asp

<210> 39

<211> 129

<212> PRT

<213> Lyssavirus sp.

<220>

<223> Mok1SAF

<400> 39

Cys Leu Asp Thr Leu Glu Thr Val Phe Met Ser Gln Ser Val Ser Phe
1 5 10 15

Arg Arg Leu Ser His Phe Arg Lys Leu Val Pro Gly Tyr Gly Lys Ala
20 25 30

Tyr Thr Ile Leu Asn Gly Ser Leu Met Glu Ala Asn Val Tyr Tyr Lys
35 40 45

Arg Val Asp Arg Trp Ala Asp Ile Leu Pro Ser Lys Gly Cys Leu Lys 50 55 60

Val Gly Gln Gln Cys Met Asp Pro Val Asn Gly Val Leu Phe Asn Gly 65 70 75 80

Ile Ile Lys Gly Pro Asp Gly Gln Ile Leu Ile Pro Glu Met Gln Ser 85 90 95

```
Glu Gln Leu Lys Gln His Met Asp Leu Leu Lys Ala Ala Val Phe Pro
                                 105
Leu Arg His Pro Leu Ile Ser Gln Glu Ala Val Phe Lys Lys Asp Gly
                             120
                                                  125
Asp
<210> 40
<211> 522 .
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Consensus
      sequence
<220>
<221> MOD RES
<222> (2)
<223> variable amino acid
<220>
<221> MOD RES
<222> (9)
<223> variable amino acid
<220>
<221> MOD_RES
<222> (13)..(15)
<223> variable amino acid
<220>
<221> MOD_RES
<222> (58)
<223> variable amino acid
<220>
<221> MOD RES
<222> (111)
<223> variable amino acid
<220>
<221> MOD RES
<222> (181)
<223> variable amino acid
<220>
<221> MOD RES
<222> (189)
<223> variable amino acid
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<220>

<221> MOD_RES <222> (262)

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<223> variable amino acid.
<220>
<221> MOD_RES
<222> (264)..(265)
<223> variable amino acid
<220>
<221> MOD RES
<222> (267)..(268)
<223> variable amino acid
<220>
<221> MOD_RES
<222> (310)
<223> variable amino acid
<220>
<221> MOD_RES
<222> (355)
<223> variable amino acid
<220>
<221> MOD RES
<222> (374)
<223> variable amino acid
<220>
<221> MOD RES
<222> (445)
<223> variable amino acid
<220>
<221> MOD RES
<222> (488)..(489)
<223> variable amino acid
<220>
<221> MOD_RES
<222> (494)
<223> variable amino acid
<220>
<221> MOD RES
<222> (518)
<223> variable amino acid
<220>
<221> MOD RES
<222> (521)
<223> variable amino acid
<400> 40
Met Xaa Leu Gln Ala Val Ile Phe Xaa Leu Leu Xaa Xaa Xaa Leu
Cys Leu Gly Lys Phe Pro Ile Tyr Thr Ile Pro Asp Lys Leu Gly Pro
```

25

20

30

Trp Ser Pro Ile Asp Ile His His Leu Ser Cys Pro Asn Asn Leu Val 35 40 45

Val Glu Asp Glu Gly Cys Thr Thr Leu Xaa Pro Phe Ser Tyr Met Glu 50 55 60

Leu Lys Val Gly Tyr Ile Thr Thr Ile Lys Val Ser Gly Phe Thr Cys
65 70 75 80

Thr Gly Val Val Thr Glu Ala Glu Thr Tyr Thr Asn Phe Val Gly Tyr 85 90 95

Val Thr Thr Phe Lys Arg Lys His Phe Arg Pro Thr Val Xaa Ala 100 105 110

Cys Arg Asp Ala Tyr Asn Trp Lys Ile Ala Gly Asp Pro Arg Tyr Glu 115 120 125

Glu Ser Leu His Asn Pro Tyr Pro Asp Ser His Trp Leu Arg Thr Val 130 135 140

Thr Thr Lys Glu Ser Leu Leu Ile Ile Ser Pro Ser Val Val Asp 145 150 155 160

Met Asp Ala Tyr Asp Lys Thr Leu His Ser Lys Met Phe Pro Asn Gly
165 170 175

Lys Cys Ser Gly Xaa Ser Pro Ser Ser Pro Phe Cys Xaa Thr Asn His 180 185 190

Asp Tyr Thr Ile Trp Met Pro Glu Asn Pro Asn Pro Gly Leu Ser Cys
195 200 205

Asp Ile Phe Thr Thr Ser Lys Gly Lys Lys Ala Thr Lys Gly Gly Arg 210 215 220

Leu Cys Gly Phe Val Asp Glu Arg Gly Leu Tyr Lys Ser Leu Lys Gly
225 230 235 240

Ala Cys Lys Leu Lys Leu Cys Gly Ile Pro Gly Leu Arg Leu Met Asp 245 250 255

Gly Ser Trp Val Ser Xaa Gln Xaa Xaa Glu Xaa Xaa Lys Trp Cys Ser 260 265 270

Pro Asp Gln Leu Val Asn Ile His Asp Phe His Ser Asp Glu Ile Glu 275 280 285

His Leu Ile Val Glu Glu Leu Val Lys Lys Arg Glu Glu Cys Leu Asp 290 295 300

Ala Leu Glu Thr Ile Xaa Thr Thr Lys Ser Ile Ser Phe Arg Arg Leu 305 310 315 320

Ser His Leu Arg Lys Leu Val Pro Gly Phe Gly Lys Ala Tyr Thr Ile 325 330 335 Ile Asn Lys Thr Leu Met Glu Ala Asp Ala His Tyr Lys Ser Val Arg 340 345 350

Glu Trp Xaa Glu Ile Ile Pro Ser Lys Gly Cys Leu Lys Ala Gly Gly 355 360 365

Arg Cys His Pro His Xaa Asn Gly Val Phe Phe Asn Gly Ile Ile Leu 370 380

Gly Pro Asp Gly His Val Leu Ile Pro Glu Met Gln Ser Ala Leu Leu 385 390 395 400

Gln Gln His Ile Glu Leu Leu Glu Ser Ser Val Ile Pro Leu Arg His 405 410 415

Pro Leu Ala Asp Pro Ser Thr Val Phe Lys Lys Asp Asp Glu Ala Glu 420 425 430

Asp Phe Val Glu Val His Leu Pro Asp Val Gln Lys Xaa Ile Ser Gly 435 440 445

Ile Asp Leu Gly Leu Pro Glu Trp Lys Arg Tyr Phe Leu Ile Gly Ala 450 455 460

Ser Ala Leu Ala Leu Leu Ala Leu Ala Ile Ile Ile Ala Val Cys Cys 465 470 475 480

Lys Arg Val Lys Arg Arg Arg Xaa Xaa Lys Pro Asn Pro Xaa Glu Leu 485 490 495

Ile Arg Lys Val Ser Val Thr Ser Gln Ser Gly Lys Val Ile Pro Ser 500 505 510

Trp Glu Ser Tyr Lys Xaa Gly Thr Xaa Gly
515 520